

0500 OIPE

CRF Errors Corrected by the STIC Systems Branch

Serial Number: 10/002,585

CRF Processing Date: 12/17/2001  
Edited by: me  
Verified by: me (STIC staff)

**ENTERED**

#2

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: \_\_\_\_\_
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: \_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file;  
☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☐ Other: \_\_\_\_\_

Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form. 2/1/95

OIPE

**RAW SEQUENCE LISTING**

DATE: 12/17/2001

PATENT APPLICATION: US/10/002,585

TIME: 20:10:14

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\12172001\J002585.raw

**SEQUENCE LISTING**

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3 (1) GENERAL INFORMATION:
5   (i) APPLICANT: Yan et al.
7   (ii) TITLE OF INVENTION: COMPOUNDS AND METHODS FOR
8       MODULATION OF CD154
10  (iii) NUMBER OF SEQUENCES: 2
12  (iv) CORRESPONDENCE ADDRESS:
13      (A) ADDRESSEE: COR THERAPEUTICS, INC.
14      (B) STREET: 256 E. Grand Ave.
15      (C) CITY: South San Francisco
16      (D) STATE: CA
17      (E) COUNTRY: USA
18      (F) ZIP: 94080
20  (v) COMPUTER READABLE FORM:
21      (A) MEDIUM TYPE: Diskette
22      (B) COMPUTER: IBM PC compatible
23      (C) OPERATING SYSTEM: PC-DOS/MS-DOS
24      (D) SOFTWARE: Patentin Release #124
26  (vi) CURRENT APPLICATION DATA:
C--> 27      (A) APPLICATION NUMBER: US/10/002,585
C--> 28      (B) FILING DATE: 30-Nov-2001
C--> 29      (C) CLASSIFICATION:
31  (vii) PRIOR APPLICATION DATA:
32      (A) APPLICATION NUMBER: US 60/289,049
33      (B) FILING DATE: 03-MAY-2001
C--> 35  (viii) ATTORNEY/AGENT INFORMATION:
36      (A) NAME: Pili Ekstrom, Carmen B.
C--> 37      (B) REGISTRATION NUMBER: 34,981
C--> 38      (C) REFERENCE/DOCKET NUMBER: COR 148
C--> 40  (ix) TELECOMMUNICATION INFORMATION:
41      (A) TELEPHONE: (650) 244-6896
42      (B) TELEFAX: (650) 246-7896
48 (2) INFORMATION FOR SEQ ID NO: 1:
49   (i) SEQUENCE CHARACTERISTICS:
50      (A) LENGTH: 35 base pairs
51      (B) TYPE: nucleic acid
52      (C) STRANDEDNESS: single
53      (D) TOPOLOGY: linear
55   (ii) MOLECULE TYPE: other nucleic acid
56      (A) DESCRIPTION: /desc = "Synthetic DNA"
58   (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
60   CGAATTCCTC TTCCATGGAA AACAGCTTTG AAATG   35
62 (2) INFORMATION FOR SEQ ID NO: 2:
64   (i) SEQUENCE CHARACTERISTICS:
65      (A) LENGTH: 29 base pairs
66      (B) TYPE: nucleic acid
67      (C) STRANDEDNESS: single

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**RAW SEQUENCE LISTING**

PATENT APPLICATION: US/10/002,585

DATE: 12/17/2001

TIME: 20:10:14

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\12172001\J002585.raw

68 (D) TOPOLOGY: linear  
70 (ii) MOLECULE TYPE: other nucleic acid  
71 (A) DESCRIPTION: /desc = "Synthetic DNA"  
73 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:  
75 GACTCTTCGA AGCTAGGATC CTAGGGTTA 29

**VERIFICATION SUMMARY**

PATENT APPLICATION: US/10/002,585

DATE: 12/17/2001

TIME: 20:10:15

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\12172001\J002585.raw

L:27 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]  
L:28 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]  
L:29 M:220 C: Keyword misspelled or invalid format, [(C) CLASSIFICATION:]  
L:29 M:220 C: Keyword misspelled or invalid format, Poss data loss, (C) CLASSIFICATION:  
L:35 M:220 C: Keyword misspelled or invalid format, [(viii) ATTORNEY/AGENT INFORMATION:]  
L:37 M:220 C: Keyword misspelled or invalid format, [(B) REGISTRATION NUMBER:]  
L:38 M:220 C: Keyword misspelled or invalid format, [(C) REFERENCE/DOCKET NUMBER:]  
L:40 M:220 C: Keyword misspelled or invalid format, [(ix) TELECOMMUNICATION INFORMATION:]